

SEQUENCE LISTING

<110> Endress, Gregory A.
Rosen, Craig A.

<120> Prostate Specific Secreted Protein

<130> PF457

<140> Unassigned

<141> 1999-03-30

<150> 60/080,311

<151> 1998-04-01

<150> 60/080,898

<151> 1998-04-07

<160> 15

<170> PatentIn Ver. 2.0

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ccagtcaca ggcgagagcc ctggg atg cac cgg cca gag gcc atg ctg ctg 172
                               Met His Arg Pro Glu Ala Met Leu Leu
                               1           5

ctg ctc acg ctt gcc ctc ctg ggg ggc ccc acc tgg gca ggg aag atg 220
Leu Leu Thr Leu Ala Leu Leu Gly Gly Pro Thr Trp Ala Gly Lys Met
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tat ggc cct gga gga ggc aag tat ttc agc acc act gaa gac tac gac 268
Tyr Gly Pro Gly Gly Gly Lys Tyr Phe Ser Thr Thr Glu Asp Tyr Asp
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cat gaa atc aca ggg ctg cgg gtg tct gta ggt ctt ctg ctg gtg aaa 316
His Glu Ile Thr Gly Leu Arg Val Ser Val Gly Leu Leu Leu Val Lys
           45           50           55

agt gtc cag gtg aaa ctt gga gac tcc tgg gac gtg aaa ctg gga gcc 364
Ser Val Gln Val Lys Leu Gly Asp Ser Trp Asp Val Lys Leu Gly Ala
           60           65           70

tta ggt ggg aat acc cag gaa gtc acc ctg cag cca ggc gaa tac atc 412
Leu Gly Gly Asn Thr Gln Glu Val Thr Leu Gln Pro Gly Glu Tyr Ile
           75           80           85

aca aaa gtc ttt gtc gcc ttc caa gct ttc ctc cgg ggt gtg gtc atg 460
Thr Lys Val Phe Val Ala Phe Gln Ala Phe Leu Arg Gly Val Val Met
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tac acc agc aag gac cgc tat ttc tat ttt ggg aag ctt gat ggc cag 508
Tyr Thr Ser Lys Asp Arg Tyr Phe Tyr Phe Gly Lys Leu Asp Gly Gln
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atc tcc tct gcc tac ccc agc caa gag ggg cag gtg ctg gtg ggc atc 556
Ile Ser Ser Ala Tyr Pro Ser Gln Glu Gly Gln Val Leu Val Gly Ile

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aat tat cca cta gag gag ccg acc act gag cca cca gtt aat ctc aca 652
 Asn Tyr Pro Leu Glu Glu Pro Thr Thr Glu Pro Pro Val Asn Leu Thr
 155 160 165

tac tca gca aac tca ccc gtg ggt cgc tag ggtggggtat ggggccatcc 702
 Tyr Ser Ala Asn Ser Pro Val Gly Arg
 170 175

gagctgaggc catctgggtg gtggtggctg atggtactgg agtaactgag tcgggacgct 762

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 35 40 45

Val Ser Val Gly Leu Leu Leu Val Lys Ser Val Gln Val Lys Leu Gly
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Asp Ser Trp Asp Val Lys Leu Gly Ala Leu Gly Gly Asn Thr Gln Glu
 65 70 75 80

Val Thr Leu Gln Pro Gly Glu Tyr Ile Thr Lys Val Phe Val Ala Phe
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Gln Ala Phe Leu Arg Gly Val Val Met Tyr Thr Ser Lys Asp Arg Tyr
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Phe Tyr Phe Gly Lys Leu Asp Gly Gln Ile Ser Ser Ala Tyr Pro Ser
 115 120 125

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 130 135 140

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Gln Glu Tyr Gly Ser Ser Gly Arg Ala Glu Ile Glu Val Lys Leu Asn			
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Pro Asp Glu Thr Val Leu Gly Phe Ser Gly Ser Phe Tyr Ile Phe Met			
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His Gln Ile Ile Ile Thr Thr Ser Gln Pro Arg Glu Leu Ile Ile Gly			
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Pro Leu Thr Gly Arg Tyr Val Tyr Thr Ser Tyr Pro Glu Asn Pro Asn			
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 gccctaact ccgcccagtt ccgcccattc tccgcccatt ggctgactaa ttttttttat 180
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